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L Holm, C Sander - Bioinformatics, 1998 - Oxford Univ Press

... Whenever there is not a match, one cannot be certain ... Wootton, JC (1990) Construction of validated, non-redundant composite protein sequence databases. ...

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The DNA sequence of human chromosome 21- ➤ fil-leibniz.de (por)

M Hattori, A Fujiyama, TD Taylor, H Watanabe ... - Nature, 2000 - nature.com ... of >0.6 and had no match to repetitive ... Alu (red) and LINE1 (blue) interspersed repeat

element densities ... The final non-redundant sequence was divided into 340 ...

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AT Castelo, W Martins, GR Gao - Bioinformatics, 2002 - Oxford Univ Press

... m is a match for a motif of length i at position k in the DNA sequence. ... to account for the appearance of these redundant repeats a master record R i is ...

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R Radtkey, L Feng, M Muralhidar, M Duhon, D ... - Nucleic Acids Research, 2000 - Oxford Univ Press

... the 1 repeat reporter, the match signal appears ... provides the opportunity for redundant interrogation of ... interrogate a slightly different sequence position of ...

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A map for **sequence** analysis of the Arabidopsis thaliana genome

M Marra, T Kucaba, M Sekhon, LD Hillier, R ... - nature genetics, 1999 - nature.com

... succeed in providing high-quality sequence-ready material ... The high redundancy achieved by the large number ... best-matching fingerprint.) The best match to clone ...

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[PDF] ... RPE/choroid for the NEIBank Project: over 6000 non-redundant transcripts, novel genes ...

G Wistow, SL Bernstein, MK Wyatt, RN Fariss, ... - Mol Vis, 2002 - molvis.org

... NEIBank Project: Over 6000 non-redundant transcripts, novel ... from lens (by), the

RPE/choroid sequence data have a ... 45% of the clusters do not match named genes ...

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... contains potential coding regions among pseudogenes and a high density of repeat ...- Full Text@IngentaConnect

M Van Geel, LJ Heather, R Lyle, JE Hewitt, ... - Genomics, 1999 - Elsevier

... clone was initially sequenced with a sixfold redundancy. ... A 50-kb sequence fragment containing FRG1 and ... predictions usually do not exactly match the boundaries ...

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RV Kantety, M La Rota, DE Matthews, ME ... - Plant Molecular Biology, 2002 - Springer ... the dataset and creates a longer consensus **sequence** by de ... singletons with a total of 4231 non-**redundant**(NR) sequences ... of 85% similarity in order to **match** as many ... Cited by 250 - Related articles - All 7 versions

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